

Enhancing Hit Identification in *Mycobacterium tuberculosis* Drug Discovery Using Dual-Event Bayesian Models

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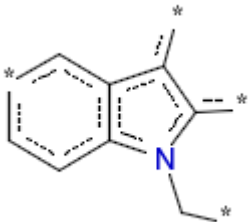
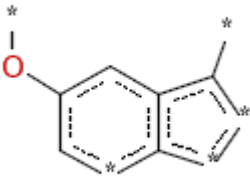
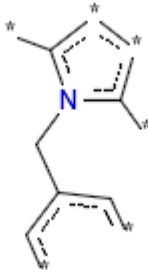
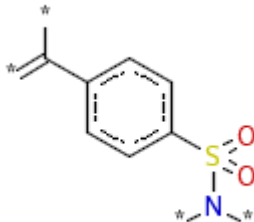
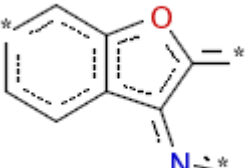
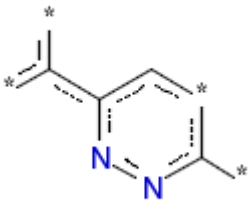
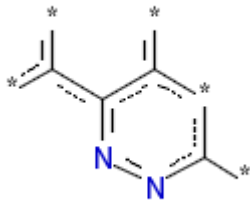
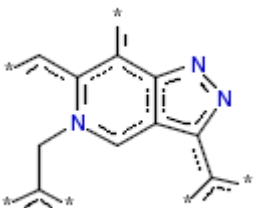
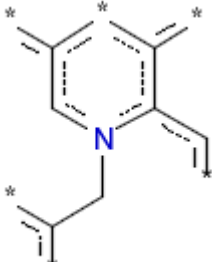
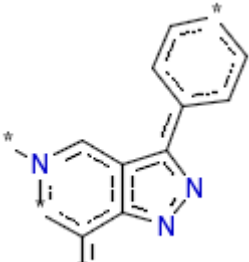
⁶Department of Medicine, Center for Emerging and Reemerging Pathogens, UMDNJ – New Jersey Medical School, 185 South Orange Avenue Newark, NJ 07103, USA.

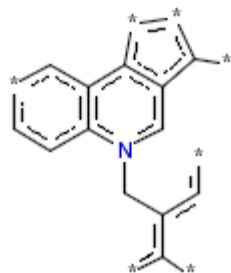
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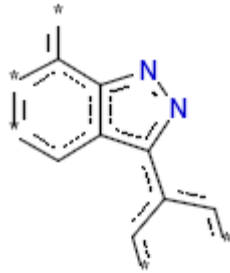
Running Head: Dual Event Bayesian Models

Figure S8. TB kinase dose response and cytotoxicity model: bad features from FCFP_6.

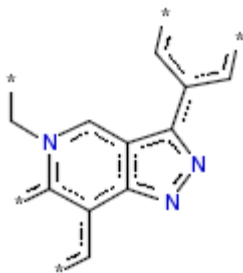
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 <p>B6: 1964502305 0 out of 37 good Bayesian Score: -1.835</p>	 <p>B7: 1087307120 0 out of 37 good Bayesian Score: -1.835</p>	 <p>B8: 1813774889 0 out of 36 good Bayesian Score: -1.812</p>	 <p>B9: -1731186150 0 out of 36 good Bayesian Score: -1.812</p>	 <p>B10: -1236091207 0 out of 36 good Bayesian Score: -1.812</p>



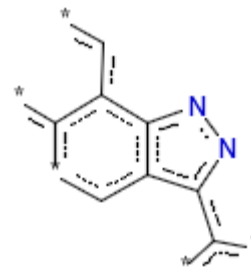
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Bayesian Score: -1.812



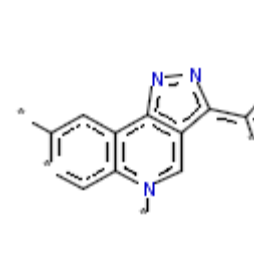
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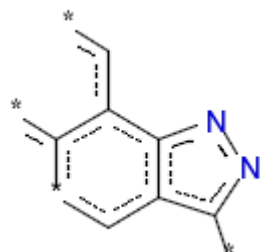
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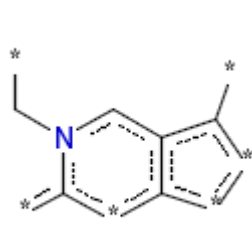
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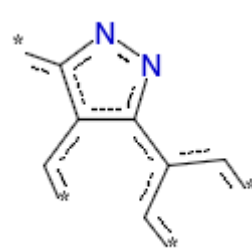
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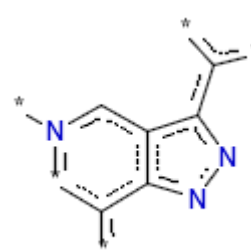
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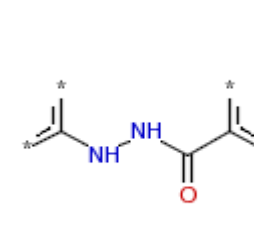
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Bayesian Score: -1.812



B18: 789148918
0 out of 36 good
Bayesian Score: -1.812



B19: -1227636663
0 out of 36 good
Bayesian Score: -1.812



B20: 1935427586
0 out of 34 good
Bayesian Score: -1.765